



#10

R0133045.txt
SEQUENCE LISTING

<110> Kane, Michael D.
Nagel, Aaron C.
Dombkowski, Alan A.

<120> COMPOSITIONS AND SYSTEMS FOR IDENTIFYING AND COMPARING EXPRESSED GENES
(mRNAs) IN EUKARYOTIC ORGANISMS

<130> 65446-87

<160> 5

<170> PatentIn version 3.1

<210> 1

<211> 61

<212> DNA

<213> Artificial Sequence

<220>

<223> This is a synthesized sequence.

<220>

<221> misc_feature

<222> (1)..(50)

<223> The poly-T tail may have between 8 and 50 residues.

<220>

<221> misc_feature

<222> (52)..(62)

<223> The sequence may have between 3 and 10 residues selected from a
or c or g or t.

<400> 1
tttttttttt tttttttttt tttttttttt tttttttttt vnnnnnnnnnn 60
n 61

<210> 2

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

```

<223> This is a synthesized sequence.
<220>
<221> misc_feature
<222> (23)..(25)
<223> n may be selected from a or c or g or t.
<400> 2
tttttttttt ttttttttt tvnnn
25
<210> 3
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> This is a synthesized sequence.
<400> 3
tttttttttt ttttttttt aaac
24
<210> 4
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> This is a synthesized sequence.
<400> 4
gtttaaaaaa aaaaaaaaaa aaaa
24
<210> 5
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> This is a synthesized sequence.
<400> 5
tttttttttt ttttttttt gggtt
25

```